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Timestamp: Thu Sep 27 12:25:44 EDT 2007

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Application No: 10798579 Version No: 3.0

Input Set:

Output Set:

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Total Warnings: 2  
Total Errors: 0  
No. of SeqIDs Defined: 68  
Actual SeqID Count: 68

Error code	Error Description
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<110> SHINOZAKI, KAZUKO  
 UMEMOTO, NAOYUKI  
 MAMIYA, KANJI  
 TOGURI, TOSHIHIRO

<120> PRODUCTION OF PLANTS HAVING IMPROVED ROOTING EFFICIENCY  
 AND VASE LIFE USING STRESS-RESISTANCE GENE

<130> 081356-0210

<140> 10798579

<141> 2004-03-12

<150> JP 2003-71082

<151> 2003-03-14

<160> 68

<170> PatentIn Ver. 3.3

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atg aac tca ttt tct gct ttt tct gaa atg ttt ggc tcc gat tac gag 166

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Ser Ser Val Ser Ser Gly Gly Asp Tyr Ile Pro Thr Leu Ala Ser Ser

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tgc ccc aag aaa ccg gcg ggt cgt aag aag ttt cgt gag act cgt cac 262

Cys Pro Lys Lys Pro Ala Gly Arg Lys Lys Phe Arg Glu Thr Arg His

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cca ata tac aga gga gtt cgt cgg aga aac tcc ggt aag tgg gtt tgt 310

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Glu Val Arg Glu Pro Asn Lys Lys Thr Arg Ile Trp Leu Gly Thr Phe

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Pro Ile Tyr Arg Gly Val Arg Arg Arg Asn Ser Gly Lys Trp Val Cys  
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 Glu Val Arg Glu Pro Asn Lys Lys Thr Arg Ile Trp Leu Gly Thr Phe  
 65 70 75 80  
 Gln Thr Ala Glu Met Ala Ala Arg Ala His Asp Val Ala Ala Leu Ala  
 85 90 95  
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 100 105 110  
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 Ala Glu Ala Ala Leu Ala Phe Gln Asp Glu Met Cys Asp Ala Thr Thr  
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 Asp His Gly Phe Asp Met Glu Glu Thr Leu Val Glu Ala Ile Tyr Thr  
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Lys Ala Asp Lys His Trp Leu Ser Glu Phe Glu His Asn Tyr Trp Ser	
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Asp Ile Leu Lys Glu Lys Glu Lys Gln Lys Glu Gln Gly Ile Val Glu	
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Thr Cys Gln Gln Gln Gln Gln Asp Ser Leu Ser Val Ala Asp Tyr Gly	
230 235 240	

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 Asp Val Asp Glu Leu Leu Arg Asp Leu Asn Gly Asp Asp Val Phe Ala  
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 Tyr Arg Pro Glu Ser Gln Gln Ser Gly Phe Asp Pro Leu Gln Ser Leu  
 295 300 305  
  
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 Asn Tyr Gly Ile Pro Pro Phe Gln Leu Glu Gly Lys Asp Gly Asn Gly  
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 Val Ser Thr Lys Lys Arg Lys Val Pro Ala Lys Gly Ser Lys Lys Gly  
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 Gly Val Arg Gln Arg Ile Trp Gly Lys Trp Val Ala Glu Ile Arg Glu

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Gly Val Glu Pro Met Tyr Cys Leu Glu Asn Gly Ala Glu Glu Met Lys		
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Arg Gly Val Lys Ala Asp Lys His Trp Leu Ser Glu Phe Glu His Asn		
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Tyr Trp Ser Asp Ile Leu Lys Glu Lys Glu Lys Gln Lys Glu Gln Gly		
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Ile Val Glu Thr Cys Gln Gln Gln Gln Gln Asp Ser Leu Ser Val Ala		
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275	280	285
Asn Gly Ser Tyr Arg Pro Glu Ser Gln Gln Ser Gly Phe Asp Pro Leu		
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Gln Ser Leu Asn Tyr Gly Ile Pro Pro Phe Gln Leu Glu Gly Lys Asp		
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Ser Ala Phe Ser Glu Met Phe Gly Ser Asp Tyr Glu Pro Gln Gly Gly  
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gat tat tgt ccg acg ttg gcc acg agt tgt ccg aag aaa ccg gcg ggc 271  
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25 30 35

cgt aag aag ttt cgt gag act cgt cac cca att tac aga gga gtt cgt 319  
Arg Lys Lys Phe Arg Glu Thr Arg His Pro Ile Tyr Arg Gly Val Arg  
40 45 50

caa aga aac tcc ggt aag tgg gtt tct gaa gtg aga gag cca aac aag 367  
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Lys Thr Arg Ile Trp Leu Gly Thr Phe Gln Thr Ala Glu Met Ala Ala  
70 75 80

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Arg Ala His Asp Val Ala Ala Leu Ala Leu Arg Gly Arg Ser Ala Cys  
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tgc gcc aag gat atc caa aaa gcg gct gct gaa gcg gcg ttg gct ttt 559  
Cys Ala Lys Asp Ile Gln Lys Ala Ala Ala Glu Ala Ala Leu Ala Phe  
120 125 130

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